SEQUENCE LISTING

SEQ ID NO: 1:

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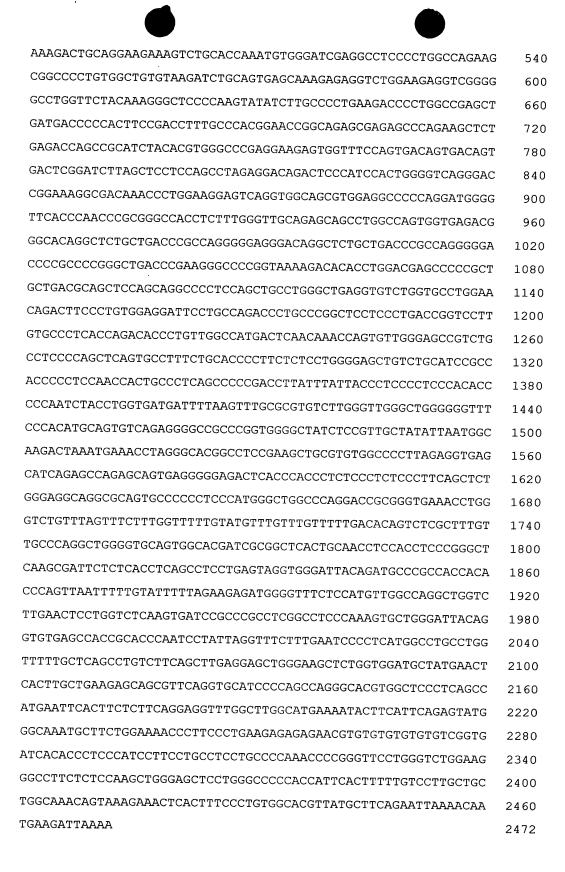


SEQ ID NO: 2:

 ${\tt MetAlaAspThrIlePheGlySerGlyAsnAspGlnTrpValCysProAsnAspArgGln}$ 20 LeuAlaLeuArgAlaLysLeuGlnThrGlyTrpSerValHisThrTyrGlnThrGluLys 40 ${ t GlnArgArgLysGlnHisLeuSerProAlaGluValGluAlaIleLeuGlnValIleGln}$ 60 ${\tt ArgAlaGluArgLeuAspValLeuGluGlnGlnArgIleGlyArgLeuValGluArgLeu}$ 80 GluThrMetArgArgAsnValMetGlyAsnGlyLeuSerGlnCysLeuLeuCysGlyGlu 100 ValLeuGlyPheLeuGlySerSerSerValPheCysLysAspCysArgLysValTrpLys 120 ${\tt ArgSerGlyAlaTrpPheTyrLysGlyLeuProLysTyrIleLeuProLeuLysThrPro}$ 140 ${\tt GlyArgAlaAspGluProGlnPheArgProTrpProThrGluProAlaGluArgGluPro}$ 160 ${\tt ArgSerSerGluThrSerArgIleTyrThrTrpAlaArgGlyArgValValSerSerAsp}$ 180 SerAspSerAspSerAspLeuSerSerSerLeuGluAspArgLeuProSerThrGly 200 ValArgAspArgLysGlyAspLysProTrpLysGluSerGlyGlySerValGluAlaPro 220 ${\tt ArgMetGlyPheThrGlnProAlaGlyHisLeuPheGlyLeuGlnSerSerLeuAlaSer}$ 240 260 ${\tt ProGlyGlyProArgProGlyLeuThrArgArgAlaProValLysAspThrProGlyArg}$ 280 AlaProAlaAlaAspAlaAlaProAlaGlyProSerSerCysLeuGly 296

SEQ ID NO: 3:

60 $\tt CTCCTCCTGGTGGGGCCTGTCTGGGTGAAGCCCCTCTGTTCCCGAGGATCGTCCCA$ 120 ACCCCCAGCCGGGTGCTCCGAGCCATGGCCGACACCATCTTCGGCAGCGGGAATGATCAG 180 TGGGTTTGCCCCAATGACCGGCAGCTTGCCCTTCGAGCCAAGCTGCAGACGGGCTGGTCC 240 GTGCACACCTACCAGACGGAGAAGCAGAGGAAGCAGCACCTCAGCCCGGCGGAGGTG 300 GAGGCCATCCTGCAGGTCATCCAGAGGGCAGAGCGGCTCGACGTCCTGGAGCAGCAGAGA 360 ATCGGGCGGCTGGAGCCGCTGAGACCATGAGGCGGAATGTGATGGGGAACGGCCTG 420 TCCCAGTGTCTGCTGCGGGGGGGGGGTGCTGGGCTTCCTGGGCAGCTCGTCGGTGTTCTGC 480



SEO ID NO: 4:

 ${\tt MetAlaAspThrIlePheGlySerGlyAsnAspGlnTrpValCysProAsnAspArgGln}$ 20 ${\tt LeuAlaLeuArgAlaLysLeuGlnThrGlyTrpSerValHisThrTyrGlnThrGluLys}$ 40 ${ t GlnArgArgLysGlnHisLeuSerProAlaGluValGluAlaIleLeuGlnValIleGln}$ 60 ${\tt ArgAlaGluArgLeuAspValLeuGluGlnGlnArgIleGlyArgLeuValGluArgLeu}$ 80 ${\tt GluThrMetArgArgAsnValMetGlyAsnGlyLeuSerGlnCysLeuLeuCysGlyGlu}$ 100 ${\tt ValLeuGlyPheLeuGlySerSerValPheCysLysAspCysArgLysLysValCys}$ 120 ${\tt ThrLysCysGlyIleGluAlaSerProGlyGlnLysArgProLeuTrpLeuCysLysIle}$ 140 ${\tt CysSerGluGlnArgGluValTrpLysArgSerGlyAlaTrpPheTyrLysGlyLeuPro}$ 160 LysTyrIleLeuProLeuLysThrProGlyArgAlaAspAspProHisPheArgProLeu 180 ${\tt ProThrGluProAlaGluArgGluProArgSerSerGluThrSerArgIleTyrThrTrp}$ 200 $\verb|AlaArgGlyArgValValSerSerAspSerAspSerAspSerAspLeuSerSerSerSer| \\$ 220 ${\tt LeuGluAspArgLeuProSerThrGlyValArgAspArgLysGlyAspLysProTrpLys}$ 240 ${\tt GluSerGlyGlySerValGluAlaProArgMetGlyPheThrGlnProAlaGlyHisLeu}$ 260 ${\tt PheGlyLeuGlnSerSerLeuAlaSerGlyGluThrGlyThrGlySerAlaAspProPro} \\$ 280 GlyGlyGlyThrGlySerAlaAspProProGlyGlyProArgProGlyLeuThrArgArg 300 ${ t AlaProValLysAspThrProGlyArgAlaProAlaAlaAspAlaAlaProAlaGlyPro}$ 320 SerSerCysLeuGly 325

SEQ ID NO: 5:

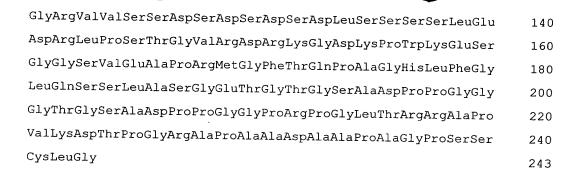
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4

AGGGACCGGAAAGGCGACAAACCCTGGAAGGAGTCAGGTGGCAGCGTGGAGGCCCCCAGG 960 ATGGGGTTCACCCAACCCGCGGGCCACCTCTTTGGGTTGCAGAGCAGCCTGGCCAGTGGT 1020 1080 GGGGGACCCCGCCCGGGCTGACCCGAAGGGCCCCGGTAAAAGACACACCTGGACGAGCC 1140 $\verb|CCCGCTGCTGACGCAGCTCCAGCTGCCTGGGCTGAGGTGTCTGGTGC| \\$ 1200 CTGGAACAGACTTCCCTGTGGAGGATTCCTGCCAGACCCTGCCCGGCTCCTCCCTGACCG 1260 $\tt GTCCTTGTGCCCTCACCAGACACCCTGTTGGCCATGACTCAACAACCAGTGTTGGGAGC$ 1320 CGTCTGCCTCCCCAGCTCAGTGCCTTTCTGCACCCCTTCTCTCTGGGGAGCTGTCTGCA 1380 1440 CACACCCCCAATCTACCTGGTGATGATTTTAAGTTTGCGCGTGTCTTGGGTTGGGCTGGG 1500 GGGTTTCCCACATGCAGTGTCAGAGGGGCCCCCGGTGGGGCTATCTCCGTTGCTATATT 1560 AATGGCAAGACTAAATGAAACCTAGGGCACGGCCTCCGAAGCTGCGTGTGGCCCCTTAGA 1620 1680 AGCTCTGGGAGGCAGGCGCAGTGCCCCCTCCCATGGGCTGGCCCAGGACCGCGGGTGAA 1740 ACCTGGGTCTGTTTAGTTTCTTTGGTTTTTGTATGTTTTGTTTTTTGACACAGTCTCG 1800 $\tt CTTTGTTGCCCAGGCTGGGGTGCAGTGGCACGATCGCGGCTCACTGCAACCTCCACCTCC$ 1860 CGGGCTCAAGCGATTCTCTCACCTCAGCCTCCTGAGTAGGTGGGATTACAGATGCCCGCC 1920 ACCACACCCAGTTAATTTTTGTATTTTTAGAAGAGATGGGGTTTCTCCATGTTGGCCAGG 1980 CTGGTCTTGAACTCCTGGTCTCAAGTGATCCGCCCGCCTCGGCCTCCCAAAGTGCTGGGA 2040 TTACAGGTGTGAGCCACCGCACCCAATCCTATTAGGTTTCTTTGAATCCCCTCATGGCCT 2100 ${\tt GCCTGGTTTTTGCTCAGCCTGTCTTCAGCTTGAGGAGCTGGGAAGCTCTGGTGGATGCTA}$ 2160 2220 2280 2340 TCGGTGATCACACCCTCCCATCCTTCCTGCCTCCTGCCCCAAACCCCGGGTTCCTGGGTC 2400 $\tt TGGAAGGGCCTTCTCCCAAGCTGGGAGCTCCTGGGCCCCCACCATTCACTTTTTGTCCT$ 2460 TGCTGCTGGCAAACAGTAAAGAAACTCACTTTCCCTGTGGCACGTTATGCTTCAGAATTA 2520 AAACAATGAAGATTAAAA 2538

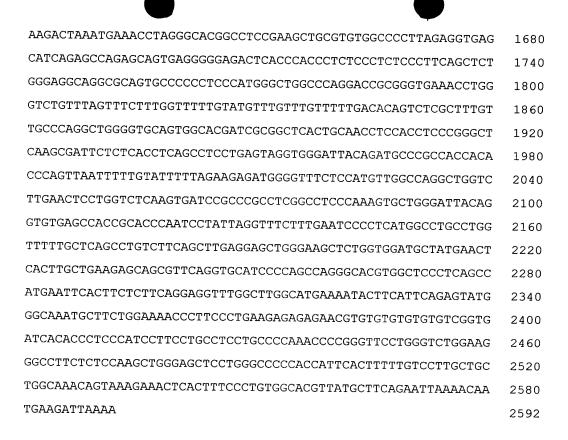
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SEQ ID NO: 7:

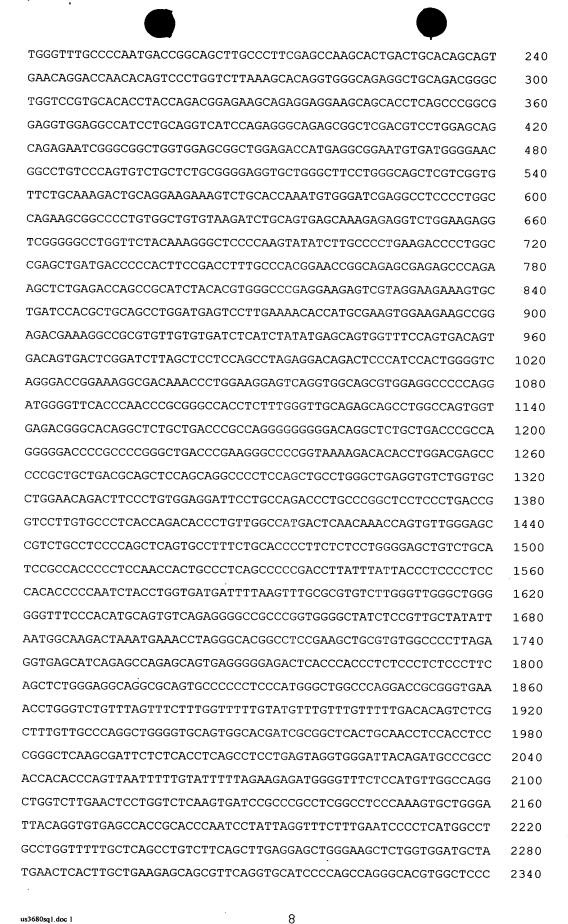
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GAGGCCATCCTGCAGGTCATCCAGAGGGCAGAGCGGCTCGACGTCCTGGAGCAGCAGAGA	360
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CCTCCCCAGCTCAGTGCCTTTCTGCACCCCTTCTCTCTGGGGAGCTGTCTGCATCCGCC	1440
ACCCCTCCAACCACTGCCCTCAGCCCCGACCTTATTTATT	1500
CCCAATCTACCTGGTGATGATTTTAAGTTTGCGCGTGTCTTGGGTTGGGCTGGGGGGTTT	1560
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SEQ ID NO: 8:

 ${\tt MetAlaAspThrIlePheGlySerGlyAsnAspGlnTrpValCysProAsnAspArgGln}$ 20 LeuAlaLeuArgAlaLysLeuGlnThrGlyTrpSerValHisThrTyrGlnThrGluLys 40 ${\tt GlnArgArgLysGlnHisLeuSerProAlaGluValGluAlaIleLeuGlnValIleGln}$ 60 ArgAlaGluArgLeuAspValLeuGluGlnGlnArgIleGlyArgLeuValGluArgLeu 80 GluThrMetArgArgAsnValMetGlyAsnGlyLeuSerGlnCysLeuLeuCysGlyGlu 100 ValLeuGlyPheLeuGlySerSerSerValPheCysLysAspCysArgLysLysValCys 120 ThrLysCysGlyIleGluAlaSerProGlyGlnLysArgProLeuTrpLeuCysLysIle 140 ${\tt CysSerGluGlnArgGluValTrpLysArgSerGlyAlaTrpPheTyrLysGlyLeuPro}\\$ 160 ${\tt LysTyrIleLeuProLeuLysThrProGlyArgAlaAspAspProHisPheArgProLeu}$ 180 ${\tt ProThrGluProAlaGluArgGluProArgSerSerGluThrSerArgIleTyrThrTrp}$ 200 AlaArgGlyArgValValGlyArgLysCys 210

SEQ ID NO: 9:





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